

GenCore version 4.5
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Result No.	Query No.	Score	Match Length	DB ID	Description
Run on:	January 3, 2002, 19:19:14 ;	Search time 4748.56 Seconds (without alignments)			
		17655.598 Million cell updates/sec			
Title:	US-09-497-82224-18				
Perfect score:	5082				
Sequence:	1 gagctctggacaaaaatttag.....acaagcaaaaaaaa 5082				
Scoring table:	IDENTITY_NUC				
	Gapop 10.0 , Gapext 1.0				
Searched:	1472140 seqs, 824858975 residues				
	Total number of hits satisfying chosen parameters:	2944280			
	Minimum DB seq length: 0				
	Maximum DB seq length: 2000000000				
Post-processing:	Minimum Match 0%				
	Maximum Match 100%				
	L <small>IST</small> ING first 45 summaries				
Database :	GenEmbl:*				
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	2 gb_ltg:*				
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	4: gb_om:*				
	5: gb_ov:*				
	6: gb_rat:*				
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ALIGNMENTS	
RESULT	1
AR153576	
LOCUS	AR153576
DEFINITION	3715 bp DNA
ACCESSION	Sequence 10 from Patent US 62235872
VERSION	AR153576
KEYWORDS	GI:15121108
SOURCE	Unknown.
ORGANISM	Unclassified.
REFERENCE	1 (bases 1 to 3715)
AUTHORS	Bredesen,D.E. and Rabizadeh,S.
TITLE	proapoptotic peptides dependence on
JOURNAL	Patent: US 6235872-A 10-22 MAY-2000
FEATURES	Location/Qualifiers
source	1 .. 3715 /organism="unknown"
BASE COUNT	841 a 1055 c 1001 g
ORIGIN	818
SUMMARIES	

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and Δ is derived by the formula $\Delta = \frac{1}{2} \sum_{i=1}^{n-1} (i - \text{Pred. No.})$.

the total

FEATURES	source	Location/Qualifiers
BASE COUNT		1 -3715 /organism="unknown"
ORIGIN	841 a	1055 c 1001 g

TITLE Proapoptotic peptides dependence polypeptides and methods of use
JOURNAL
Patent: US 6235872-A 10-22-MAY-2001;

Search completed: January 3, 2002, 21:35:15

